**Amendments to the Specification:**

Please replace the paragraph beginning at page 15, line 12 with the following paragraph:

The post-expression truncation of the wild-type enzyme produces a mixture of truncated enzyme molecules with a molecular weight ranging from 27 kDa to 37.5 kDa. However, there is a dominant species, referred as TF-glucanase, that has a molecular weight of approximately 27.7 kDa, that is, about 10 kDa smaller than that of the full-length enzyme. This dominant TF-glucanase is formed after 10 to 14 days post-expression incubation in the LB medium at 25 °C, and is stable and active even if when the incubation time is extended for up to 45 days at 25 °C. The sequence of this dominant TF-glucanase is presented in FIG 2, which suggests that TF-glucanase is produced when approximately 80 amino acid residues are removed from the C-terminus of the wild-type enzyme. On the other hand, the PCR-generated truncated enzyme, i.e., PCR-TF-glucanase, has a molecular weight of 29.7 kDa, and shares the same amino acid sequence with TF-glucanase except that PCR-TF-glucanase has 19 extra amino acid residues at the C-terminus, see FIG 3. Five P-X-S-S-S-S repeats and a basic terminal domain (BTD) located in the C-terminal portion of the wild-type enzyme (SEQ ID NO:11) are absent from either TF-glucanase or PCR-TF-glucanase. The symbol P represents ~~poline~~ proline, S presents serine, and X represents an uncharged residue, such as alanine ~~Alanine~~, proline ~~Proline~~, or glutamine ~~Glutamine~~.